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D-81927 München (DE)(71) Applicant: BOEHRINGER MANNHEIM GMBH
68298 Mannheim (DE)**(54) Immunoregulatory protein LST-1**

(57) The invention concerns a new immunoregulatory protein LST-1, nucleic acid sequences coding for this protein, a process for the isolation of this protein, as well as its use for the production of a therapeutic agent. The DNA and protein sequences are shown in SEQ ID NO:1 to 4.

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Description

This invention relates to a new leukocyte specific and immunoregulatory protein (LST-1) and to the corresponding protein produced by recombinant techniques as well as nucleic acids which code for proteins with LST-1 activity, methods of use and of production.

Cytokines are proteins with a molecular weight of less than 50 kD, which mediate the exchange of autocrine, paracrine or endocrine signals between the cellular components of tissues or between different tissues. The cytokines identified so far include growth factors, interleukins and interferons and act on cells in many systems of the body: the hematopoietic system, the immune system, the nervous system, the skeletal system, connective tissues, and probably most other tissues and organs of the body (for reference see A. Thompson ed. (1991), The Cytokine Handbook, London, Academic Press and A. Miyajima et al., Annual Reviews Immunol. 10 (1992) 295-331). Examples of cytokines are EGF, NGF, PDGF, FGF, IL-1 to IL-7, GM-CSF, G-CSF, MCSF, IFN, TNF- α , TGF- α and - β .

More than 100 cytokines have been identified so far, but there is a need for further new cytokines which might be important for potential health care advances.

Summary of the invention

Accordingly it is an object of the present invention to provide a new cytokine showing new biological properties.

It is a further object of the present invention to provide the new protein using recombinant DNA molecules capable of expressing said protein in order that the binding protein will be more readily available. These and other objects of the invention have been accomplished by providing a purified protein according to the invention, selected from a group consisting of a protein which is at least 85% homologous to the amino acid sequence SEQ ID NO:3 and fragments thereof, wherein said protein is capable of binding to an antibody specific for said protein or its cell surface receptor on leukocytes.

Description of specific embodiments

The invention comprises especially novel therapeutic compositions comprising recombinant proteins produced using nucleic acid sequences encoding proteins with LST-1 activity. The LST-1 cDNA which can be used for the production of the recombinant protein was initially isolated from U937 cells stimulated with IFN- α . RT-PCR cloning of mRNA from these cells resulted in a cDNA clone designated pLST-1 having a length of 636 bp. Analysis of the LST-1 cDNA reveals that LST-1 consists of six exons. These exons are designated exons 1A (bp 48-162), 1B (bp 544-652), 2 (bp 1044-1162), 3 (bp 1475-1567), 4 (bp 1775-1797) and 5 (bp 2325-2709). In exon 5, after position 2345, there is an internal 5' donor splice site. By alternative splicing, thus, two isoforms of the LST-1 protein can be formed, which accordingly have a length of either 104 or 97 amino acids (cf. SEQ ID NOS: 3 and 4). The human pLST-1 cDNA clone contains an extended 5' region encoding stop codons.

The invention is based on a new cytokine-like protein (denoted LST-1 protein, or LST-1, in the following) whose production is stimulated in U937 cells by IFN- γ by a factor of more than 1000, which binds to the surface of leukocytes and which

- a) is coded by the DNA sequence shown in SEQ ID NO:2 for the mature protein or by the genomic sequence shown in SEQ ID NO:1.
- b) is coded by DNA sequences which hybridize under stringent conditions with the DNA sequences shown in SEQ ID NO:1 or 2 or fragments of the DNA sequences in the DNA region which codes for the mature protein, or
- c) is coded by DNA sequences which, if there was no degeneracy of the genetic code, would hybridize with the sequences defined in a) or b) and code for a polypeptide with amino acid sequence.

The protein can be defined by its DNA sequence and by the amino acid sequence derived therefrom. The LST-1 protein can occur in natural allelic variations which differ from individual to individual. Such variations of the amino acids are usually amino acid substitutions. However, they may also be deletions, insertions or additions of amino acids to the total sequence. The LST-1 protein according to the invention - depending, both in respect of the extent and type, on the cell and cell type in which it is expressed - can be in glycosylated or non-glycosylated form. The LST-1 proteins according to SEQ ID NO:3 and 4 are preferred.

LST-1 mRNA is expressed constitutively in T cells, macrophages U937 and at low levels in human tonsilla, lung and liver, which may be due to the lymphocytes and macrophages present in these tissues. The protein shows an immunoregulatory activity. Transcription can be strongly enhanced by IFN- γ in U937 cells. Upon stimulation of the Jurkat T cell line (ATCC TIB 152) with TPA (12-o-tetradecanoylphorbol-13-acetate) only a short induction of LST-1 mRNA was observed. An increase of mRNA expression starts after four hours with a peak after 8 hours of stimulation and further decrease after 12 hours of stimulation.

"Immunoregulatory activity" means that the protein modulates directly or non-directly the cooperation of T-cells with macrophages.

The binding to the surface of leucocytes is preferably estimated in vitro. Such methods are known in the state of the art.

5 The term "hybridize under stringent conditions" means that two nucleic acid fragments are capable of hybridization to one another under standard hybridization conditions described in Sambrook et al., "Expression of cloned genes in E. coli" in Molecular Cloning: A laboratory manual (1989) Cold Spring Harbor Laboratory Press, New York USA, 9.47 - 9.62 and 11.45 - 11.61.

10 More specifically, stringent conditions as used herein refer to hybridization in 1 mol/l NaCl, 1 % SDS and 10 % dextran sulphate. This is followed by two washes of the filter at room temperature of 5 minutes in 2 x SSC and one final wash for 30 minutes. This final wash may be at 0.5 x SSC, 0.1 % SDS, more preferably at 0.2 x SSC, 0.1 % SDS and most preferably at 0.1 x SSC, 0.1 % SDS, final wash taking place at 65°C. Those of ordinary skill in the art will recognize that other conditions will afford the same degrees of stringency and are encompassed by the phraseology "under stringent conditions" and are encompassed herein.

15 LST-1 is a protein which is active in its glycosylated or unglycosylated form. The unglycosylated form can be produced by recombinant technology in prokaryotic cells.

"Proteins with LST-1 activity" means also proteins with minor amino acid variations but with substantially the same LST1 activity. Substantially the same means that the activities are of the same biological properties and preferably at least 85% homology in amino acid sequence. More preferably, the amino acid sequences are at least 90% identical.

20 LST-1 can be purified from U937 cells by affinity chromatography using a monoclonal antibody against LST-1. It is also preferred to use other known protein purification techniques, including immunoprecipitation, gel filtration, ion exchange, chromatography, chromatofocusing, isoelectric focusing, selective precipitation, electrophoresis, and the like. Fraction isolated during purification procedures can be analyzed for the presence of LST-1 activity by using LST-1 specific antibodies.

25 The protein according to the invention can also be produced by recombinant means. Non-glycosylated LST-1 protein is obtained when it is produced recombinantly in prokaryotes. With the aid of the nucleic acid sequences provided by the invention it is possible to search for the LST-1 gene or its variants in genomes of any desired cells (e.g. apart from human cells, also in cells of other mammals), to identify these and to isolate the desired gene coding for the LST-1 protein. Such processes and suitable hybridization conditions are known to a person skilled in the art and are described, for example, by Sambrook, J., et al., "Expression of cloned genes in E. coli" in Molecular Cloning: A laboratory manual (1989) Cold Spring Harbor Laboratory Press, New York, USA, and B.D. Hames, S.G. Higgins, Nucleic acid hybridisation - a practical approach (1985) IRL Press, Oxford, England. In this case the standard protocols described in these publications are usually used for the experiments.

30 The use of recombinant DNA technology enables the production of numerous LST-1 protein derivatives. Such derivatives can, for example, be modified in individual or several amino acids by substitution, deletion or addition. The derivatization can, for example, be carried out by means of site directed mutagenesis. Such variations can be easily carried out by a person skilled in the art (J. Sambrook, B.D. Hames, loc. cit.). It merely has to be ensured that the characteristic properties of the LST-1 protein (inhibition of the aforementioned cell lines) are preserved. The invention therefore in addition concerns a LST-1 protein which is a product of a prokaryotic or eukaryotic expression of an exogenous DNA.

40 The invention concerns a nucleic acid molecule for use in securing expression in a prokaryotic or eukaryotic host cell of a LST-1 protein and is selected from the group comprising

- a) DNA sequences shown in SEQ ID NO:1 and 2 or the complementary sequences,
- b) nucleic acid sequences which hybridize under stringent conditions with one of the sequences from a),
- 45 c) nucleic acid sequences which, if there was no degeneracy of the genetic code, would hybridize with one of the sequences stated in a) or b).

With the aid of such nucleic acids coding for a LST-1 protein, the protein according to the invention can be obtained in a reproducible manner and in large amounts. For expression in prokaryotic or eukaryotic organisms, such as prokaryotic host cells or eukaryotic host cells, the nucleic acid is integrated into suitable expression vectors, according to methods familiar to a person skilled in the art. Such an expression vector preferably contains a regulatable/inducible promoter. These recombinant vectors are then introduced for the expression into suitable host cells such as, e.g., E. coli as a prokaryotic host cell or *Saccharomyces cerevisiae*, Teratocarcinoma cell line PA-1 sc 9117 (Büttner et al., Mol. Cell. Biol. 11 (1991) 3573-3583), insect cells, CHO or COS cells as eukaryotic host cells and the transformed or transduced host cells are cultured under conditions which allow an expression of the heterologous gene. The isolation of the protein can be carried out according to known methods from the host cell or from the culture supernatant of the host cell. Such methods are described for example by Ausubel L, Frederick M., Current Protocols in Mol. Biol. (1992), John Wiley and Sons. . . York. Also in vitro reactivation of the protein may be necessary.

In addition the invention concerns a process for obtaining a LST-1 protein by isolation of the culture supernatant of

the U937 cell line by means of a gel chromatographic separation and purification of a fraction.

The detection of transformed or transduced host cells which recombinantly produce the LST-1 protein and the purification of the protein are preferably carried out by means of antibodies which bind to this protein. Such antibodies can be obtained in a simple manner according to known methods by using the protein according to the invention as an antigen or an immunogen.

The invention therefore in addition concerns the use of the protein with LST-1 activity according to the invention for the production of antibodies which bind to this protein.

Anti-LST-1 antibodies are produced by immunization and appropriate vertebrate host with purified LST-1 or polypeptide derivatives of LST-1, preferably with an adjuvant. Said techniques are well-known in the literature and are described, for example, by Harlow and Lane eds., *Antibodies: A laboratory manual* (1988), Cold Spring Harbor Laboratories Press.

For this, animals which are usually used for this purpose, such as in particular, sheep, rabbits or mice, are immunized with the protein according to the invention and subsequently the antiserum is isolated from the immunized animals according to known methods or spleen cells of the immunized animals are fused with immortalized cells, such as e.g. myeloma cells, according to the method of Köhler and Milstein (*Nature* 256 (1975) 495-497). Those cells which produce a monoclonal antibody against the LST-1 protein are selected from the hybridoma cells obtained in this way and cloned. The monoclonal or polyclonal antibodies obtained in this way can be bound to a support material, such as e.g. cellulose, for an immunoabsorptive purification of the melanoma-inhibiting protein. Furthermore, antibodies of this kind can be used for the detection of the LST-1 protein in samples, such as e.g. cut tissue or body fluids.

The invention therefore additionally concerns antibodies against the LST-1 protein which are obtainable by immunizing an animal with a LST-1 protein and isolating the antibodies from the serum or spleen cells of the immunized animals.

It has in addition turned out that the LST-1 protein has an immunoregulatory activity.

The invention in addition concerns the use of a protein according to the invention for the production of a therapeutic agent which can be used in tumor therapy or as an immunoregulating agent.

The protein according to the invention is processed, if desired together with the usually used auxiliary agents, fillers and/or additives, in a pharmaceutical formulation for the said therapeutic applications.

The invention therefore in addition concerns a therapeutic composition containing a LST-1 protein according to the invention and if desired together with the auxiliary agents, fillers and/or additives that are usually used.

The invention further concerns the use of sequences of the LST-1 gene, preferably nucleic acid molecules coding for a protein having LST-1 activity, or activating polynucleotides from the 5' untranslated region, in gene therapy, and in particular, for the production of medicaments for gene therapy.

Gene therapy of somatic cells can be accomplished by using, e.g., retroviral vectors, other viral vectors, or by non-viral gene transfer (for clarity cf. T. Friedmann, *Science* 244 (1989) 1275; Morgan 1993, RAC DATA MANAGEMENT REPORT, June 1993).

Vector systems suitable for gene therapy are, for instance, retrovirus (Mulligan, R.C. (1991) in Nobel Symposium 8: *Etiology of human disease at the DNA level* (Lindsten, J. and Patterson Editors) 143-189, Raven Press), adeno associated virus (McLughlin, J. *Virol.* 62 (1988), 1963), vaccinia virus (Moss et al., *Ann. Rev. Immunol.* 5 (1987) 305), bovine papilloma virus (Rasmussen et al., *Methods Enzymol.* 139 (1987) 642) or viruses from the group of the herpes viruses such as Epstein Barr virus (Margolskee et al., *Mol. Cell. Biol.* 8 (1988) 2937) or herpes simplex virus.

There are also known non-viral delivery systems. For this, usually "nude" nucleic acid, preferably DNA, is used, or nucleic acid together with an auxiliary such as, e.g., transfer reagents (liposomes, dendromers, polylysine-transferrine-conjugates (Felgner et al., *Proc. Natl. Acad. Sci. USA* 84 (1987) 7413).

Another preferred method of gene therapy is based on homologous recombination. In this, either the gene coding for the LST-1 protein can be inserted in one or more copies into the genome of somatic cells and/or the LST-1 gene endogenously present in the cells can be modulated, preferably activated.

Methods of homologous recombination are described, e.g., in Kucherlapati, *Proc. in Nucl. Acids Res. and Mol. Biol.* 36 (1989) 301; Thomas et al., *Cell* 44 (1986) 419-428; Thomas and Capecchi, *Cell* 51 (1987) 503-512; Doetschman et al., *Proc. Natl. Acad. Sci. USA* 85 (1988) 8583-8587 and Doetschman et al., *Nature* 330 (1987) 576-578. In these methods, a portion of DNA to be integrated at a specific site in the genome (gene fragment of LST-1) is bound to a targeting DNA. The targeting DNA is a DNA which is complementary (homologous) to a region (preferably within or proximal to the LST-1 gene) of the genomic DNA. When two homologous portions of a single-stranded DNA (e.g. the targeting DNA and the genomic DNA) are in close proximity to one another they will hybridize and form a double-stranded helix. Then the LST-1 gene fragment and the targeting DNA can be integrated into the genome by means of occurrence of recombination. This homologous recombination can be carried out both *in vitro* and *in vivo* (in the patient).

Preferably, there is used a DNA which codes for a protein having LST-1 activity, a fragment which inhibits LST-1 expression (knock-out sequence) or a fragment capable of activating, after integration of the genome of a cell, expression, in this cell, of a protein having LST-1 activity. Such a fragment may be, for example, a promoter and/or enhancer region which is heterologous to the corresponding LST-1 region or which, after integration into the LST-1 gene, activates

the actually silent or to a little extent expressed LST-1 gene transcriptionally and/or translationally.

Thus, by means of this DNA, one or more LST-1 genes are newly introduced into the target cell, or the essentially transcriptionally silent gene in the genome of a mammalian cell is activated in such fashion that the mammalian cell is enabled to produce endogenous LST-1 protein. To this end, a DNA construct is inserted into the genome by homologous recombination, the DNA construct comprising the following: a DNA regulatory element capable of stimulating expression of this gene if operatively linked thereto; and one or more DNA target segments which are homologous to a region in this genome, which region is within or proximal to this gene. This construct is inserted into the genome of the mammalian cell in such fashion that the regulatory segment is operatively linked to the gene which codes for the protein having LST-1 activity. Preferably, the construct further comprises amplifying sequences, especially if genes coding for proteins with LST-1 activity are inserted into the cell.

For the introduction of LST-1 genes into the target cells, the construct comprises a regulatory element, one or more LST-1 genes and one or more target segments. The target segments are chosen in such a way that they hybridize with an appropriate region of the genome, whereby, after homologous recombination, the inserted exogenous LST-1 genes are expressed.

There are known a large number of processes by which homologous recombination can be initiated. Preferably, homologous recombination takes place during DNA replication or mitosis of the cells. A DNA of this kind can be used for the production of an agent for therapeutic treatment of tumors or for the production of homologous or heterologous LST-1 protein in a host organism.

It is possible to provide a test on the basis of the nucleic acid sequences of the LST-1 protein provided by the invention which can be used to detect nucleic acids which code for LST-1 proteins. Such a test can for example be carried out in cells or cell lysates and by means of nucleic acid diagnostics. In this case the sample to be examined is brought into contact with a probe which would hybridize with the nucleic acid sequence coding for the LST-1 protein. A hybridization between the probe and nucleic acids from the sample indicates the presence of expressed LST-1 proteins. Such methods are known to a person skilled in the art and are for example described in WO 89/06698, EP-A 0 200 362, USP 2915082, EP-A 0 063 879, EP-A 0 173 251, EP-A 0 128 018. In a preferred embodiment of the invention, the nucleic acid of the sample which codes for a LST-1 protein is amplified before testing, e.g. by the well-known PCR technique. A derivatized (labelled) nucleic acid probe is usually used in the field of nucleic acid diagnostics. This probe is brought into contact with a carrier-bound denatured DNA or RNA from the sample and in this process the temperature, ionic strength, pH value and other buffer conditions are selected in such a way that - depending on the length of the nucleic acid sample and the resulting melting temperature of the expected hybrid - the labelled DNA or RNA can bind to homologous DNA or RNA (hybridization, see also Southern, E.M., J. Mol. Biol. 98 (1975), 503 - 517; Wahl, G.M. et al., Proc. Natl. Acad. Sci. USA 76 (1979), 3683 - 3687). Suitable carriers are membranes or carrier materials based on nitrocellulose (e.g. Schleicher and Schüll, BA 85, Amersham Hybond C.), reinforced or bound nitrocellulose in a powder form or nylon membranes derivatized with various functional groups (e.g. nitro group) (e.g. Schleicher and Schüll, Nytran; NEN, Gene Screen; Amersham Hybond M.; Pall Biodyne).

The hybridized DNA or RNA is then detected by incubating the carrier, after thorough washing and saturation to prevent unspecific binding, with an antibody or antibody fragment. The antibody or antibody fragment is directed towards the substance incorporated into the nucleic acid probe during the derivatization. The antibody is in turn labelled. It is, however, also possible to use a directly labelled DNA. After incubation with the antibodies, it is washed again in order to only detect specifically bound antibody conjugates. The determination is then carried out via the label of the antibody or antibody fragment according to well-known methods.

The detection of the LST-1 expression can be carried out for example as:

- in situ hybridization with immobilized whole cells using immobilized tissue smears and isolated metaphase chromosomes,
- colony hybridization (cells) and plaque hybridization (phages and viruses),
- Northern hybridization (RNA detection),
- serum analysis (e.g. cell type analysis of cells in serum by slot-blot analysis),
- after amplification (e.g. PCR technique).

The invention therefore includes a method for the detection of nucleic acids which code for a LST-1 protein which is characterized in that the sample to be examined is incubated with a nucleic acid probe which is selected from the group comprising

- 55 a) the DNA sequences shown in SEQ ID NOS:1 and 2 or a complementary sequence to these,
- b) nucleic acids which hybridize under stringent conditions with one of the sequences from a), the nucleic acid probe is incubated with the nucleic acid from the sample and the hybridization of the nucleic acid in the sample and nucleic acid probe is detected, if desired, via a further binding partner.

Thus, LST-1 is a valuable prognostic marker in tumor diagnostics (metastasis, progress) and an activity marker for cell proliferation, especially for T cell leukemic cells.

Plasmid pLST-1 which contains the LST-1 gene was deposited by Boehringer Mannheim GmbH at the Deutsche Sammlung von Mikroorganismen und Zellkulturen GmbH, Mascheroder Weg 1b, D-38124 Braunschweig on 24 May 5 1995 and assigned the number DSM 10011.

The invention is elucidated in more detail by the sequence protocols in conjunction with the following examples and figures. In this case

SEQ ID NO 1 denotes the nucleotide sequence of the LST-1 genomic region. A polymorphic Pvu II restriction site 10 is at position 2841. The TATA box, the IFN- γ -activated site (FcgR1), and the interferon-stimulated gene factor-2 responsive element (ISGF-2) are starting at positions 279, 1509, and 1814, respectively. The DNA sequence has been submitted to GenBank databank (accession number U00921).

SEQ ID NO 2 denotes protein and cDNA of LST-1.

SEQ ID NO 3 denotes LST-1 protein sequence (104 amino acids).

SEQ ID NO 4 denotes LST-1 protein sequence (isoform 97 amino acids)

Abbreviations: BAT, HLA-B-associated transcript; MHC, major histocompatibility complex; LST-1, Leucocyte Specific Transcript - 1; pLst1, LST-1 cDNA clone; MER, medium reiteration frequency repeat; TNF- α , tumor necrosis factor - α ; TNFA, gene coding for TNF- α ; TNFB, gene coding for tumor necrosis factor - β ; LTB, gene coding for lymphotoxin - β .

Example 1

Isolation of the LST-1 genomic region

The human B cell line CAH was established from a homozygous individual typed HLA-A3, -Bw47, -Cw6, and -DR7. 30 Genomic DNA was used to generate the cosmid library cah in the vector pTCF as described previously (E.H. Weiss et al., Immunobiology 170 (1985) 367-380). Several cosmids were isolated by hybridization to a TNFA probe. A subclone derived from the cosmid cah5 containing the region upstream of the LTB gene was obtained by Hind III endonuclease restriction of cah5. Religation of the digest resulted in a truncated clone (cah5dH3) of 18 kb length. Three adjacent Pst I fragments of 1.5 kb, 0.8 kb, and 3.1 kb referred to as Pst6, Pst4, and Pst11, respectively, were isolated and subcloned 35 into the vector pUC19.

Example 2

Determination of the nucleotide sequence

The subclones Pst4, Pst6, and Pst11 were sequenced in both directions with the dideoxynucleotide chain termination method using either Sequenase 2.0 (USB, Braunschweig, Germany) or T7 polymerase (Pharmacia, Freiburg, Germany). The genomic organization of these fragments was determined by direct sequencing of cah5dH3 across the Pst I restriction sites, employing oligonucleotides. This confirmed the order and excluded the possibility that any small fragment is situated between the restriction fragments Pst6, Pst4, and Pst11. 45

Example 3

Cell culture and stimulation experiments

Non-adherent human Jurkat T cells, the human histiocytic cell line U937, and the human monocytic cell line MonoMac6 were grown in RPMI 1640 medium. The adherent A431 and HeLa epithelial carcinoma lines were grown in Dulbecco's modified Eagle's medium (DMEM). For both media 10% heat inactivated fetal calf serum, 1% penicillin/streptomycin and 1% L-glutamine (all purchased from Gibco-BRL, Berlin, Germany) were used as a supplement. All 50 cells were cultured at 5% CO₂ and 37°C in a humidified atmosphere. Stimulation of Jurkat cells was carried out with 50 ng/ml 12-O-tetradecanoylphorbol-13-acetate (TPA) (Sigma, Deisenhofen, Germany) and 5 µg/ml phytohemagglutinin (PHA) (Sigma). The human monocytic cell line MonoMac6 was stimulated with 1 mg/ml lipopolysaccharides (prepared from Salmonella minnesota; Sigma) for 4 hours. The histiocytic cell line U937 was stimulated with 200 U/ml interferon- γ for 48 hours.

Example 4**Cloning and characterization of the LST-1 cDNA**

5 Reverse transcription of RNA, followed by the polymerase chain reaction (RT-PCR) of mRNA isolated from U937 cells stimulated with interferon- γ , resulted in a cDNA clone, designated pLst1 of 636 bp length. This is in good agreement with the size of mRNA detected by Northern blot of about 800 bp length, due to an expected length of the poly(A) tail of about 200 bp. The cDNA sequence of pLst1 was identical to the coding regions of the genomic sequence cloned from a different human cell line. Sequence analysis revealed three regions which are well conserved between the
 10 human LST-1 and the mouse B144 transcript, and could be assigned to exon 2-4 of the human gene.

Like the other genes in the TNF region, LST-1 consists of four exons and three introns. Its transcription orientation is the same as for TNFA and TNFB, but opposite to the neighboured LTB gene. This result does not agree with the orientation of the mouse B144 gene as marked in the organization of the H-2D^b region published by J.M. Wroblewski et al., Immunogenetics 32 (1990) 200-204. The human pLst1 cDNA clone contains an extended 5' region encoding stop codons in all three reading frames and can therefore be considered to be full length. The LST-1 cDNA potentially encodes a transmembrane protein.
 15

Example 5**Expression of LST-1**

LST-1 mRNA is expressed constitutively in T cells, macrophages, U937, and at low levels in human tonsilla, lung, and liver, which may be due to lymphocytes and macrophages present in these tissues. In epithelial cell lines such as HeLa and A431 cells, no LST-1 transcripts were detectable (data not shown). This expression pattern prompted us to
 25 name the corresponding gene Leucocyte Specific Transcript-1 (LST-1). Transcription levels can be strongly enhanced in U937 cells by stimulation with interferon- γ , but not by lipopolysaccharides in the macrophage cell line MonoMac6. Stimulation of the Jurkat T cell line with TPA (12-O-tetradecanoylphorbol-13-acetate) induced transient expression of LST-1 mRNA above the constitutive level. Following induction, the mRNA expression level increased after 4 hours with a peak at 8 hours and decreased after 24 hours of stimulation.

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Weiss, E.H., et al., Immunobiology 170 (1985) 367-380

WO 89/06598

s Wroblewski, J.M., et al., Immunogenetics 32 (1990) 200-204

Sequence Listing**10 (1) GENERAL INFORMATION:****(i) APPLICANT:**

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20 **(ii) TITLE OF INVENTION: New immunoregulatory protein LST-1****(iii) NUMBER OF SEQUENCES: 4****(iv) COMPUTER READABLE FORM:**

- (A) MEDIUM TYPE: Floppy disk
- (B) COMPUTER: IBM PC compatible
- (C) OPERATING SYSTEM: PC-DOS/MS-DOS
- (D) SOFTWARE: PatentIn Release #1.0, Version #1.30B (EPO)

30 (2) INFORMATION FOR SEQ ID NO: 1:**(i) SEQUENCE CHARACTERISTICS:**

- (A) LENGTH: 5581 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)**40 (ix) FEATURE:**

- (A) NAME/KEY: exon
- (B) LOCATION: 48..162

45 (ix) FEATURE:

- (A) NAME/KEY: exon
- (B) LOCATION: 544..652

50 (ix) FEATURE:

- (A) NAME/KEY: exon
- (B) LOCATION: 1044..1162

(ix) FEATURE:

- (A) NAME/KEY: exon
- (B) LOCATION: 1475..1567

55 (ix) FEATURE:

- (A) NAME/KEY: exon
- (B) LOCATION: 1775..1797

(ix) FEATURE:

(A) NAME/KEY: exon
(B) LOCATION:2325..2709

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

1 GAGCCGAACCTCCTA ACAATGCTGG GGAGGAACCC AGCCTGGGG AGAAGTTAAA 60
GCCAGAGGAG GGGCAGGAAT GTCTGAGGTG GCAACACTTC TCTTCAGCCA GACAGCACTG 120
GCCAGTTGG AGTCTGTCCA TCCTGCAGGC CACAAGCTCT GGGTAAGCTG GGAATGGGCA 180
15 GGGACCTTGG TGGAAGGATG GTCACACCCC AGAGTGGGGT GAAGCTAAGA TGAGGGGAGG 240
GAGAGTATGG GTTGAGTTT CCCTGGGCCG TCGAGGAATC CTCTGAGTCT CTGCTCCCCA 300
20 AAGAAATTAA AGACAATTCA TTTCTGTGCC CACGGCCCTT ATGGCTCCAC CTGCACCTTC 360
GCTCCCCACC CCCAGAAATT CCTCTTAAAC CCAGAAGGTC CCAGTTCCA GACCCTAGTC 420
AGTATATCTG GCTCTGGGGT GAAGAGAACG GCCCCCTCTT CACCCCTAAA CAGGAACCAG 480
25 TGGTTGGAGG GGAGGAAGTG CCTGAGGGGA AGTTATGGGG CCCCAGATAAC TCCTCCATGC 540
CCCACTTCAG CCCTAGCAGC ATCTGCCTGT GGGAAAGCAGC TCTCCACACC AGCCAAGGGG 600
GCCCCCACAC TCCCGCGCTG CTCTGCGGCT CAGGGAGCAG CCCACCTGCT GGGTGTGCTG 660
30 ATATCACCT CCCTTCTTCC CCCCAGTGCC CACACCCACC CAGGCCAGG CTCCCTCCCC 720
TCCATCATCC CCTTACCAGC ACCTAGAACCC ATCCAGGGCT GAAAAGTCCC CTCCAAACCA 780
CGTGGTCAGC CCAGGGCAGA GGAAAGGGCT GGGCTCTGGA GTTGGGCAGA GCTGGCCTTA 840
35 AACCCCAGCT CCACCTTTCT GGGATGGGTG ACCTAGTAAA GTCCAGGCTT GAATCTCGGG 900
TCTTTACTTG GGCAACGGGC ACCATGATAAC CCTATGTTCT GGGGATTAGC AGTGAGGAAT 960
40 GGAAAGTGCC CAGCTCGGGT TGGCACATAA GGGAGGCTCC CCAGCCTGGG AACGATTATA 1020
ACAGAGGGCC CCTCACTTCA CAGATGAGGA ACTTGAGGCA AGTCACCAGC CCCTGATCAT 1080
TTCGCCTAAA AGAGCAAGGA CTAGAGTTCC TGACCTCCAG GCCAGTCCCT GATCCCTGAC 1140
45 CTAATGTTAT CGCGGAATGA TGGTAAGTAA AGTGTCTCTT GCATCTGCAT AGAGAGAGTC 1200
CTGGGAGCTT AGGAAGTGAT GGGGAACAGT GATGTATGCA GCTTGGACAG GCCTCTGGGG 1260
ACAGCTGGTA CAGGAGGGAA AGGGACCTCA CGGGAGGCC AGAAACCTGG TAAGAGGTGA 1320
50 GGTATTAAGG TCTGGGATGG AGAAGCTCTG AGGGTATATT TTTCTGCCTC TAAAACGTGTT 1380
GGAGAGGGAA TCTGAGAAAG CTGCAACCAA CCAGGAGGCT GGGGTACGCT GGAGAAGGAA 1440

TGGGCTTCCT AACCTTGAGC CCTCTTCCCT GAAGATATAT GTATCTACGG GGGCCTGGGG 1500
 5 CTGGGCGGGC TCCTGCTTCT GGCACTGGTC CTTCTGTCCG CCTGCCTGTG TTGGCTGCAT 1560
 CGAAGAGGTG AGCGCTGCAC TCCCTCCCTC CCCCTGCAGC AGTCCCCCT GTGCCAAC 1620
 CCCCACACGC TTTCCACTG CTTTCCAGA ACACTGCCTG GCCCTGGAGC CACTGGGAAG 1680
 10 CCAACAGGGG AGTCCACGCC TGCTGGTGGG GGGAGCCGG GAGGCGGGAG AAGCACAAAG 1740
 GGTGGCTGT GTTGAGCTTC TTCTTTCTT CCAGTAAAGA GGCTGGAGAG GAGCTGGGTG 1800
 AGTCTGGGGA CAGGGAAAGGG GGAGGGCAAG AGAGATCCTG AGTGGGTGAG TGGGGAGAAG 1860
 15 CATGGCTGAG CGCTGAGAGG AGGGTTGGGG ACCGGAGACA AGGAGAGAGA AAGTAGGAGC 1920
 ATGAGAGAGG CAGAGAAAAT CGAGGCAAAA GAGAAAGAGA AAATGAGACA GAAACCAAGA 1980
 GAAAAAGTGA GACAGAGGAT AGGAGAGACA GGGAGAAAAT GAGAGTGAGA GAGACACAAA 2040
 20 GAGAAGAGCA ATGAAAGAGA GAGAGAGAGA GAGGCTCCAG AACCAAGGCAC AGTGGCTCAC 2100
 GTCTGTCATT CCACTATCGC AAGGCTGAGG CAGGAAGATA GCTTGAGCTC AGGGGTTGAA 2160
 GACAATCCTG GACAACATAG TGGACTCTGT CTCCAAAGAA AAAAGAGAGA GAGAGAGAGA 2220
 25 GAGAGAGAGA GGGAGAGAGA GAGAGAGAGA GGGAGAGAGA TAAGAAAGGC TGGAGGTGGG 2280
 AGCAGAACTC ACAGGGAAAGG ATCTGACGCA TCGCCTCCCA TCAGCACCTT CTGTCCTGGT 2340
 30 CCCAGGCCA GGGCTCTCA GAGCAGGAAC TCCACTATGC ATCTCTGCAG AGGCTGCCAG 2400
 TGCCCAGCAG TGAGGGACCT GACCTCAGGG GCAGAGACAA GAGAGGCACC AAGGAGGATC 2460
 CAAGAGCTGA CTATGCCTGC ATTGCTGAGA ACAAAACCCAC CTGAGCACCC CAGACACCTT 2520
 35 CCTCAACCCA GGCGGTGGA CAGGGTCCCC CTGTGGTCCA GCCAGTAAAA ACCATGGTCC 2580
 CCCCACCTCT GTGTCTCAGT CCTCTCAGTC CATCTCGAGC CTCCGTTCAA ATTGATCATC 2640
 ATCAAAACTT ATGTGGCTTT TTGACCTTG AATAGGGAAT TTTTAAATT TTTTAAAT 2700
 40 TAAAATAAAA AAAACACATG GCTCACCCCT CCACCCACTC TGGGGTCAA TAGTAATTG 2760
 TTGGGTGAAT GACAGTGTTC AGGGACCCAA GCTCCCTAA CAGCCAGAAG AGGGTATGTG 2820
 TGGGCCCTGGC AGGAAAGGGC AGTTGCCAAG GAGGAGTCAT ATCTGATCCT TCCCATTCT 2880
 45 CAGGACAATC AGGCTCAGCC TCCTGGACT GGGGAAGCA GATGTGCTGA GCTCCACAT 2940
 GGTGGTGGGA GGGCGCTGG GACCACAGCC GGCAGCTGCC TTCTTGGACC TTTCCAGGTC 3000
 50 AGACCTGGTG GAAGGGAAAG TTCAGAGTTG GGGGAATCCG GAGAGAGTAG ATTTGGCATE 3060
 TGGAGAATGG AGAAGAAAAC ACTTGAGACT CATGAGGAGT TAGTGGTGGG GCAGATTTAT 3120

TGGGGTCTTT TGAAGAGGAC TAGGGACATC TGGGCTCTGG AATCACTCCT CGGGGCCCAT 3180
 5 CTGAGGAGTG GCAGTGTGTT CCCATGTGAC AGTGGCCTGG TCAGAGAGAG GACAGGAGCT 3240
 GCTCAGTGTGTT GCAGTCCCGA GGCTCTCCTC TTCCCTGGTCT CTGTCCTCCC TCCTCCACT 3300
 CTCTTACTGC CCCTCCCACATC CCGTCCACTA TTGCCCCCTGG CTCCATTACT CACATTTGCC 3360
 10 CTGGTAATAG ACGGTGTGTC CCACGGCCAC AGAGAGAAAG CTGACAGCAT AGAATCCAGC 3420
 CCGAAGAGGA GGACTGTACC AGCCCCTAG CTGAGGATGT TCTGCATGGG GCAATGGAGA 3480
 CGGGGGTTGG GGAAGAAGTG CACACAGGCT CAGGGAGGGAGGGACAC CACCTCCAG 3540
 15 CCTGCCTCCC AAGGACATTG CCTCTTGGC CTCCAGCCAG GAGGAGACAC CACCTCCAG 3600
 CATCTCACCT TTCTCCACCA CCAGCCGAGT CCCATTCCCT GTCCCGACAC CAAGGCCAG 3660
 CACCTCCACT CTGCACACGT AGATGCCTGG CGTCATGCC TCGCACGTCC CGGATGTGCA 3720
 20 GCTCAGCCTG GTGGTCATGG AGGAAACGGG AAGAACGAGA TGGAGGCGGC CCCTGAACCTC 3780
 TGGGGTTCCA TTCCACCT CCTCCCTGG AACCAACCTCA TCTCGGAACC ACGTGACGGA 3840
 25 GCCAATGGCC AGTCTCCCTT GGCTGGCATT GAAGGAGCAG GGCAGGAAGG CAGAGGATCC 3900
 TTCCAGGGTA CGAATCTCAG GGGGCTGGGA CACCCAGAGA GCACAGGAAT CCTGGGGGGA 3960
 AAAGGGAGAC CCAGAGAAAC ACCTCCCCAG TTATTCCAAA GAGAAAAGAC AACAGAGCTT 4020
 30 GGAGTAGAAC ATCCCAGCTT TCTCCAGGCA TAGGGTGCAT GGGAAATAGAT ACTTTGGGGC 4080
 CTCATTAAC CCTTCCCTCT TAACCAATCT GATTCTTAA CATTGCTTAT TAAATCATT 4140
 TTCGGCTGGG TGCAGTCGCA CGCCTGTAAT CCCAGCACTT TGGGAGGCCG AGGTGGGCCG 4200
 35 ATCACCAGGT CAGGAGATCG AGACCACCTT GGCAACATG GTGAAACCCC GTCTCTACTA 4260
 AAAAAATACA AAAATTAGCC GGGCATGGTG GTGTGCACCT GTAATCCCAG CTACTCGGGA 4320
 GGCTGAGGCA GGAGAATCGC TTGAACCCGG GAGGCAGAGG TTGCAGTGAG CCAAGATTGC 4380
 40 GCCATTGCAC TCCAGCCTGG GCGACAAAGC AAGACTCCAT CTCAAAAAT AAAAAATAA 4440
 AATCATTCTT CAAATTCTTC CTATACCAAC TCTCACTCTC ACCCTCTGCC ATCATTCTCC 4500
 45 AGCCAGTTCA GTAGTAACCT GTCTAGCTGA AATGTAAACC ATCATGGTGA ATTAAGCTC 4560
 ATTAATGAAT TGCAGCTGCC TAGTTAACTA ATATCACTCA TTATATTATG CAGGTATTAT 4620
 TTTAGTACAA ATGGCATTGT ACAGTAAGCC ATCCTTCCCTC TTTTCTTTT TTCTTTTTT 4680
 50 GAGATGGGGT CTTGCTCTGT TGCCCAAGGCT GGAATGCAGT GGTGCAATCT TGGCTCACTG 4740

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CAAACTCCGT CCCCTGGGTT CAAGCGATCC TGGTGCCTCA GCCTCCAAG TAGCTGGAC 4800
 5 TACAGGCACC CACCAACCACG ACTGGCTAAT TTTGTATTT TCAGTCGAGA CAGGGTTCC 4860
 ACCATCTGGT CTCAAACTCC TGACCTCAAG TGATCCACCC ACCTCGGACC AGGCTGGCTC 4920
 10 AAACTCCTGA TCTCAAGTGA TCCACCTGCC TCGGGCTCCC AAAGTGCCAC CCAGCCACTC 4980
 TTGGTTTCG TTAAAGAAAG TAACTAATTA AATCTCCAGG TGAAGACGTG GCCTTAATTG 5040
 GTTGAGAGATTG CTATTTAACCGTCCATGTT GATGAATTAA ACCAAATATT AAAATCCCTG 5100
 15 ATTAATTAT CTACTTAGGG AAATTTACAA GTCATTCTAT TTCAGTGGTT CTCAAACTTG 5160
 AGTGTGTATG GAAATTACCT GGAGCATCTG CTAGAACAGA TTCCCTGGCC TACCCCCCGA 5220
 20 GTTTTGACT CAGTAGGTCT GGAGTGGGGC CTAAGAATTG TTCTAGGTT CCCAGAAATC 5280
 CACATTTGA GAACTCCTGC ATTTAGTTAA TAATATGCCT GATAGTTAAG GTCTCTCAGT 5340
 25 TCATTAAGGG CAGTTTCGGC CGGGTGCAGT TCGCACGCCT ATAATCCCAA CACTTGGGA 5400
 GGCCAAGGCG AGTGGATCAC CTGAGGTCAAG GAGTTTGAGA CCAGCCTGGC CAACATGGTG 5460
 30 AACACTCGTC TCTACTAAAA ATACACAAGT TAGCCAGCAG TAATGCCATG CACCTGTAAT 5520
 CCTAGCTACT TGGGAGGCTG AGACAGGAGA ATCATTTTA CCCAGGAGGT GGAGGCTGCA 5580
 G 5581

30 (2) INFORMATION FOR SEQ ID NO: 2:

(i) SEQUENCE CHARACTERISTICS:
 35 (A) LENGTH: 312 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

40 (ix) FEATURE:
 (A) NAME/KEY: CDS
 (B) LOCATION: 1..312

45 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

ATG TTA TCG CGG AAT GAT GAT ATA TGT ATC TAC GGG GGC CTG GGG CTG	48
Met Leu Ser Arg Asn Asp Asp Ile Cys Ile Tyr Gly Gly Leu Gly Leu	
1 5 10 15	
50 GGC GGG CTC CTG CTT CTG GCA GTG GTC CTT CTG TCC GCC TGC CTG TGT	96
Gly Gly Leu Leu Leu Ala Val Val Leu Leu Ser Ala Cys Leu Cys	
20 25 30	

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TGG CTG CAT CGA AGA GTA AAG AGG CTG GAG AGG AGC TGG CAC CTT CTG 144
 Trp Leu His Arg Arg Val Lys Arg Leu Glu Arg Ser Trp His Leu Leu
 35 40 45

5 TCC TGG TCC CAG GCC CAG GGC TCC TCA GAG CAG GAA CTC CAC TAT GCA 192
 Ser Trp Ser Gln Ala Gln Gly Ser Ser Glu Gln Glu Leu His Tyr Ala
 50 55 60

10 TCT CTG CAG AGG CTG CCA GTG CCC AGC AGT GAG GGA CCT GAC CTC AGG 240
 Ser Leu Gln Arg Leu Pro Val Pro Ser Ser Glu Gly Pro Asp Leu Arg
 65 70 75 80

15 GGC AGA GAC AAG AGA GGC ACC AAG GAG GAT CCA AGA GCT GAC TAT GCC 288
 Gly Arg Asp Lys Arg Gly Thr Lys Glu Asp Pro Arg Ala Asp Tyr Ala
 85 90 95

20 TGC ATT GCT GAG AAC AAA CCC ACC 312
 Cys Ile Ala Glu Asn Lys Pro Thr
 100

(2) INFORMATION FOR SEQ ID NO: 3:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 104 amino acids
- (B) TYPE: amino acid
- (C) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

Met Leu Ser Arg Asn Asp Asp Ile Cys Ile Tyr Gly Gly Leu Gly Leu
 1 5 10 15

35 Gly Gly Leu Leu Leu Ala Val Val Leu Leu Ser Ala Cys Leu Cys
 20 25 30

Trp Leu His Arg Arg Val Lys Arg Leu Glu Arg Ser Trp His Leu Leu
 35 40 45

40 Ser Trp Ser Gln Ala Gln Gly Ser Ser Glu Gln Glu Leu His Tyr Ala
 50 55 60

45 Ser Leu Gln Arg Leu Pro Val Pro Ser Ser Glu Gly Pro Asp Leu Arg
 65 70 75 80

Gly Arg Asp Lys Arg Gly Thr Lys Glu Asp Pro Arg Ala Asp Tyr Ala
 85 90 95

50 Cys Ile Ala Glu Asn Lys Pro Thr
 100

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(2) INFORMATION FOR SEQ ID NO: 4:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 97 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS:
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

Met	Leu	Ser	Arg	Asn	Asp	Asp	Ile	Cys	Ile	Tyr	Gly	Gly	Leu	Gly	Leu
1									10					15	

Gly	Gly	Leu	Leu	Leu	Leu	Ala	Val	Val	Leu	Leu	Ser	Ala	Cys	Leu	Cys
									25				30		

Trp	Leu	His	Arg	Arg	Val	Lys	Arg	Leu	Glu	Arg	Ser	Trp	Ala	Gln	Gly
								35	40				45		

Ser	Ser	Glu	Gln	Glu	Leu	His	Tyr	Ala	Ser	Leu	Gln	Arg	Leu	Pro	Val
								50	55			60			

Pro	Ser	Ser	Glu	Gly	Pro	Asp	Leu	Arg	Gly	Arg	Asp	Lys	Arg	Gly	Thr
							65	70		75		80			

Lys	Glu	Asp	Pro	Arg	Ala	Asp	Tyr	Ala	Cys	Ile	Ala	Glu	Asn	Lys	Pro
								85	90			95			

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Claims

1. Immunoregulatory protein whose production is stimulated in U937 cell lines by IFN- γ by a factor of more than 1000, which binds on the surface of leukocytes and which
 - a) is coded by the DNA sequence shown in SEQ ID NO:2 for the mature protein or by the genomic sequence shown in SEQ ID NO:1, or
 - b) is coded by DNA sequences which hybridize with the DNA sequences shown in SEQ ID NO:1 or 2 or fragments of these DNA sequences in the DNA region which codes for the mature protein.
2. Protein as claimed in claim 1, which has a size of about 97-104 amino acids and is obtainable from the cell culture supernatant of the cell line U937 after stimulation of the cell line by IFN- γ and purification by means of gel chromatography and reverse phase HPLC.
3. Protein as claimed in claim 1 or 2, wherein
 - a) it is a product of a prokaryotic or eukaryotic expression of an exogenous DNA,
 - b) it is coded by the DNA sequence shown in SEQ ID NO:2 for the mature protein or for the protein with an N-terminal pre sequence, or by the genomic sequence shown in SEQ ID NO:1,
 - c) it is coded by DNA sequences which hybridize with the DNA sequences shown in SEQ ID NO: 1 or 2 or fragments of these DNA sequences in the DNA region which codes for the mature protein, or
 - d) is coded by DNA sequences which, if there was no degeneracy of the genetic code, would hybridize with the sequences defined in b) to c) and code for a polypeptide with the same amino acid sequence.

e) and binds on the surface of leukocytes.

4. Protein as claimed in claims 1 to 3, with the amino acid sequence of SEQ ID NO:3 or 4.
5. Nucleic acid molecule for use in securing expression in a prokaryotic or eukaryotic host cell of a protein as claimed in one of the claims 1 to 4, wherein the sequence of the nucleic acid molecule is selected from the group comprising
 - a) DNA sequences shown in SEQ ID NO:1 or 2 or the complementary DNA sequences,
 - b) nucleic acid sequences which hybridize with one of the sequences from a),
 - c) nucleic acid sequences which, if there was no degeneracy of the genetic code, would hybridize with one of the sequences stated in a) or b).
10. DNA having the sequence shown in SEQ ID NO:1.
15. DNA having the sequence shown in SEQ ID NO:2.
8. Recombinant expression vector containing a nucleic acid which codes for a protein as claimed in claims 1 to 4 and which expresses the protein coding nucleic acid in a transformed microorganism or a transformed eukaryotic cell.
20. A prokaryotic or eukaryotic host cell which is transfected with a DNA which codes for a protein as claimed in claims 1 to 4 and which can produce the said protein.
10. The host cell as claimed in claim 9, which is E. coli or a mammalian cell line.
25. 11. The use of a nucleic acid molecule which codes for a protein as claimed in claims 1 to 4, for transfection of a prokaryotic or eukaryotic organism.
12. Process for obtaining a protein which binds on the surface of leukocytes by isolation from the culture supernatant of the cell line U937 stimulated with IFN- γ , gel chromatographic separation and purifying a fraction of the supernatant that corresponds to a protein of about 97-104 amino acids.
30. 13. Process for the recombinant production of a protein which binds on the surface of leukocytes by expression of a DNA as claimed in one of claims 5 to 7 in a suitable host cell and isolation of the protein from the host cell or the culture supernatant of the host cell.
35. 14. Use of a protein which binds on the surface of leukocytes as claimed in one of claims 1 to 4 as an antigen or immunogen for the production of antibodies which bind to this protein.
40. 15. Antibody against a protein according to claims 1 to 4, which is obtainable by immunizing an animal with a protein as claimed in one of claims 1 to 4 and isolating the antibodies from the serum or spleen cells of the immunized animals.
16. Use of a protein as claimed in claims 1 to 4, for the production of a therapeutic agent, which can be used as an immunoregulatory agent.
45. 17. Therapeutic composition containing a protein as claimed in one of claims 1 to 4, if desired together with pharmaceutical auxiliary substances, fillers and/or additives.
18. Method for the detection of nucleic acids that code for a protein according to claims 1 to 4, wherein the sample to be examined is incubated with a nucleic acid probe which is selected from the group comprising
 - a) DNA with sequences shown in SEQ ID NO:1 or 2 or their complementary sequences,
 - b) nucleic acids which hybridize under stringent conditions with one of the sequences from a).
50. 55. the nucleic acid probe is incubated with the nucleic acid of the sample and the hybridization of the nucleic acid of the sample and the nucleic acid probe is detected, if desired via a further binding partner.
19. Method as claimed in claim 18, wherein the nucleic acid to be detected is amplified before detection.

20. Process for the production of a protein according to claims 1 to 4, comprising the steps of expressing in a mammalian cell the endogenous gene for this protein by inserting a DNA construct into the genome of the cell by homologous recombination, said DNA construct comprising a DNA regulatory element capable of stimulating the expression of said gene if operatively linked thereto, and further comprising one or more DNA target segments homologous to a region in this genome, which region is within or proximal to said gene; culturing the cells and recovering said protein from the cells or the cell culture supernatant.

10 21. A process for the production of a protein as claimed in claims 1 to 4, comprising the steps of expressing in a mammalian cell at least one exogenous gene which codes for this protein by inserting a DNA construct into the genome of the cell by homologous recombination, said DNA construct comprising a DNA regulatory element capable of stimulating the expression of said gene if operatively linked thereto, a DNA element which codes for said protein and further comprising one or more DNA target segments homologous to a region in this genome, culturing the cells and recovering the LST-1 protein from the cells or cell culture supernatant.

15 22. The use of a DNA construct which can be introduced into the genome of mammalian cells by homologous recombination, said DNA construct comprising

- a DNA regulatory element capable of modulating the expression of a gene if operatively linked thereto, said gene coding for a protein according to claims 1 to 4,
- and one or more DNA target segments homologous to a region in this genome, which region is within or proximal to said gene,

20 and wherein said construct can be inserted into said genome of the mammalian cell in such fashion that the regulatory element is operatively linked to the gene coding for said protein,
25 for preparing a tumor therapeutic agent or an immunoregulatory agent.

25 23. The use of a DNA construct which can be introduced into the genome of mammalian cells by homologous recombination, said DNA construct comprising

- a DNA regulatory element capable of modulating the expression of a gene if operatively linked thereto, said gene coding for a protein according to claims 1 to 4,
- and one or more DNA target segments homologous to a region in this genome where the insertion of the construct is appropriate,

30 and wherein said construct can be inserted into said genome of the mammalian cell in such fashion that the regulatory segment modulates the expression of said gene in said mammalian cell.

35 24. Plasmid DSM 10011

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EUROPEAN SEARCH REPORT

Application Number
EP 95 10 9511

DOCUMENTS CONSIDERED TO BE RELEVANT			CLASSIFICATION OF THE APPLICATION (Int.Cl.)
Category	Citation of document with indication, where appropriate, of relevant passages	Relevant to claim	
X	EMBL PRIMATE DATABANK Accession Number u00921 04-01-94 * the whole document * ---	1, 3-7	C12N15/12 C07K14/52 A61K48/00 A61K39/395 A61K38/00 C12Q1/68
X	EMBL PRIMATE DATABANK Accession number x67841 26-08-92 * the whole document * ---	1, 3-7	
Y	EP-A-0 315 950 (OTSUKA PHARMA CO LTD) 17 May 1989 * page 4, line 26 - page 5, line 32 * ---	8-13, 20-24	
Y	WO-A-87 06942 (SCRIPPS CLINIC RES) 19 November 1987 * claims 1-12 * ---	2, 22, 23	
Y	WO-A-90 13300 (BIOGEN INC) 15 November 1990 * abstract; claims 129, 130 * ---	8-24	
Y	HUM IMMUNOL, JAN 1995, 42 (1) P9-14, UNITED STATES, DE BAEY A ET AL 'Pvu II polymorphism in the primate homologue of the mouse B144 (LST-1). A novel marker gene within the tumor necrosis factor region.' * the whole document * ---	8-24	C07K C12N A61K
X	IMMUNOBIOLOGY, 191 (2-3). 1994. 149., HOLZINGER I ET AL 'LST-1, a novel gene in the human TNF region' * the whole document * -----	1-7	
The present search report has been drawn up for all claims			
Place of search	Date of completion of the search	Examiner	
BERLIN	22 November 1995	Gurdjian, D	
CATEGORY OF CITED DOCUMENTS			
X : particularly relevant if taken alone	T : theory or principle underlying the invention		
Y : particularly relevant if combined with another document of the same category	E : earlier patent document, but published on, or after the filing date		
A : technological background	D : document cited in the application		
O : non-written disclosure	L : document cited for other reasons		
P : intermediate document	& : member of the same patent family - corresponding document		